

PCT

RAW SEQUENCE LISTING DATE: 03/22/2006
PATENT APPLICATION: US/10/535,635 TIME: 15:24:37

Input Set : A:\sequence listing text.DOC
Output Set: N:\CRF4\03222006\J535635.raw

```
1 <110> APPLICANT: INCYTE CORPORATION; TRAN, Uyen K.;
             RICHARDSON, Thomas W.; BECHA, Shanya D.;
     3
             ELLIOTT, Vicki S.; SWARNAKAR, Anita;
             LEE, Soo Yeun; RAMKUMAR, Jayalaxmi;
             WANG, Jonathan T.; CHIEN, David;
     5
     6
             MURAGE, Jaji; GERA, Mili;
             MARQUIS, Joseph P.; CHAWLA, Narinder K.;
             NAKAMURA, Lisa; KABLE, Amy E.
    10 <120> TITLE OF INVENTION: IMMUNE RESPONSE-ASSOCIATED PROTEINS
    12 <130> FILE REFERENCE: PF-1629 PCT
C--> 14 <140> CURRENT APPLICATION NUMBER: US/10/535,635
C--> 15 <141> CURRENT FILING DATE: 2005-05-20
    17 <150> PRIOR APPLICATION NUMBER: US 60/429,442
    18 <151> PRIOR FILING DATE: 2002-11-26
    20 <150> PRIOR APPLICATION NUMBER: US 60/429,839
    21 <151> PRIOR FILING DATE: 2002-11-27
                                                            Does Not Comply
    23 <150> PRIOR APPLICATION NUMBER: US 60/439,946
    24 <151> PRIOR FILING DATE: 2003-01-13
                                                            Corrected Diskette Needed
    26 <150> PRIOR APPLICATION NUMBER: US 60/446,182
    27 <151> PRIOR FILING DATE: 2003-02-07
    29 <160> NUMBER OF SEQ ID NOS: 64
    30 <170> SOFTWARE: PERL Program
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ERRORED SEQUENCES

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2316 <210> SEQ ID NO: 64
2317 <211> LENGTH: 1566
2318 <212> TYPE: DNA
2319 <213> ORGANISM: Homo sapiens
2321 <220> FEATURE:
2322 <221> NAME/KEY: misc_feature
2323 <223> OTHER INFORMATION: Incyte ID No: 7514003CB1
2325 <400> SEQUENCE: 64
2326 tagggagact ctgaggetet gttgagaate atgetttgga ggeageteat etattggeaa 60
2327 ctgctggctt tgtttttcct ccctttttgc ctgtgtcaag atgaatacat ggagtctcca 120
2328 caaaccggag gactaccccc agactgcagt aagtgttgtc atggagacta cagctttcga 180
2329 qqctaccaaq qccccctqq qccaccqqqc cctcctqqca ttccaqqaaa ccatqqaaac 240
2330 aatggcaaca atggagccac tggtcatgaa ggagccaaag attgcattca tggcttctct 300
2331 ggcaacccac ttcagcaatc agaacagtgg gattatcttc agcagtgttg agaccaacat 360
2332 tggaaacttc tttgatgtca tgactggtag atttggggcc ccagtatcag gtgtgtattt 420
2333 cttcaccttc agcatgatga agcatgagga tgttgaggaa gtgtatgtgt accttatgca 480
2334 caatggcaac acagtettea geatgtacag etatgaaatg aagggcaaat cagatacate 540
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2335 cagcaatcat gctgtgctga agctagccaa aggggatgag gtttggctgc gaatgggcaa 600
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    2338 tttgttacga tctgaggaac attaaagttg agggttttac attgctgtat tcaaaaaatt 780
    2339 attggttgca atgttgttca cgctacaggt acaccaataa tgttggacaa ttcaggggct 840
    2340 caqaaqaatc aaccacaaaa tagtcttctc agatgacctt gactaatata ctcagcatct 900
    2341 ttatcactct ttccttggca cctaaaagat aattctcctc tgacgcaggt tggaaatatt 960
    2342 tttttctatc acagaagtca tttgcaaaga attttgacta ctctgctttt aatttaatac 1020
    2343 caqttttcaq qaacccctqa aqttttaagt tcattattct ttataacatt tgagagaatc 1080
    2344 agatgtagtg atatgacagg gctggggcaa gaacaggggc actagctgcc ttattagcta 1140
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    2347 tgcttgacca gtaatgtggt tgtaattttg tgtatgttcc cccacatcgc ccccaacttc 1320
    2348 gqatgtgggg tcaggaggtt gaggttcact attaacaaat gtcataaata tctcatagag 1380
    2349 qtacaqtqcc aatagatatt caaatgttgc atgttgacca gagggatttt atatctgaag 1440
    2350 aacatacact attaataaat accttagaga aagattttga cctggcttta gataaaactg 1500
    2351 tqqcaagaaa aatgtaatga qcaatatatg gaaataaaca cacctttgtt aaagataaaa 1560
                                                                        1566
    2352 aaaaaa
W--> 2356 PF-1629 PCT
E--> 2364 1/37
E--> 2365 wash 1559424.1
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VERIFICATION SUMMARY

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L:14 M:270 C: Current Application Number differs, Replaced Current Application Number

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:2356 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2

L:2364 M:254 E: No. of Bases conflict, LENGTH:Input:37 Counted:1567 SEQ:64

L:2364 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2

M:254 Repeated in SeqNo=64

L:2365 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:9

L:2365 M:112 C: (48) String data converted to lower case,

L:2365 M:252 E: No. of Seq. differs, <211> LENGTH:Input:1566 Found:1573 SEQ:64